Figure 1: Predicted protein sequence of mGy12 (SEQ ID NO:1)

- 1 MSSKTASTNS IAQARRTVQQ LRLEASIERI KVSKASADLM SYCEEHARSD
- 51 PLLMGIPTSE NPFKDKKTCI IL*

Figure 2: cDNA sequence of mGy12 variant 1 (SEQ ID NO:2)

CTAGAATTCA GCGGCCGCTG AATTCTAGGC GACGACGGCG AAGAGTGAGT GCCAAGGTTC ATATGGGAAG GACTTTGGGG TGAGCATCTT CTCTATTTCC AGCTGGCTTT TCTGATTTTC AGAAAGAAGA CTCATCAAAG ATGTCCAGCA 101 AGACGGCAAG CACCAACAGC ATAGCCCAAG CCAGGAGAAC TGTGCAGCAG 1.51 CTGAGATTGG AAGCCTCCAT CGAAAGAATA AAGGTCTCAA AAGCATCAGC 201 AGACCTGATG TCATACTGTG AGGAGCATGC CCGGAGCGAC CCCCTGCTGA 251 TGGGCATACC GACCTCAGAA AACCCGTTCA AGGATAAGAA GACCTGCATC 301 ATCTTATAGT GGACCAGGAA GCGCCCCTTG CCTCTTAACG CAAACCACAG 351 CAGCAACCTG AAGGGATTCC TTCAGCTTAC CTGGTAACCA CAGCTAGTAA 401 CTAAAACACC CTTCTCTCGG AATAATAGAC CCTGAAGTCT CTCTTTTTCA 451 AGTTGTCCTT TCTTCACCCT TTACTGATTT AATACAGAAT AACAATCTTA 501 TTTTCTATTT GATAACTATG GTATCATATT GGGTTACTGT ATAAGGAAAA 551 TGGCAGGGGA GTTGTGGGAA GCTTGTCTTT ACAAAATATA ATTGATTAAG 601 ATATGTCAAG ACCTACATTG TCTAAGCACC GGCAAATTAA AATGTCGAGA 651 701 ATCACTTCAG TCAAAAACCT TTATATTCTG TTCTTAATAA TGTTTGTGCC AACCTATATC CCATGTAAGG GATCTGGGGA GGAGGCATGT GTCTACAACC 751 ATACCTTTTT GCACTATGGG CACTAACCAC CCTGAAACTT CCTGCGGTAG 801 851 CTCCCTCCCT TCAGAGTTAC ATCATTATCC TGACTCTGTG TAGGTAAATT TCCGTGAAAT TTTTGTACAA AAAAAAGGTA ATGAAAGAAC GTTGCAAAGA 901 TCATCTGCAT TATAATGAGT TGATGCTGTT CTCACTCCTC TCTTGGAATT 951 GTGCTGGCCC CTTAGTCTAC AATAAACTGT GCCAATTAAA AACCTAAGGC 1001 TAAAACTGAA AGCCCTTTGA TGGGGTCTTA ACTCATATCA GTCATTTGGG 1051 CTTCTCTGAT CCTGAGGCTA AGAAAGGGGA AGAGACCCTC AGGAGGCAGC 1101 TTCCACTCCA GGGCTCTTGA TCTCTGCTGG ATTGGGGGTG GCCACCTCAG 1151 AAACTTCCAC CCTCATGACT GGAATGGAAG AGGGGACCGA GAGCCTCACA 1201

1251 ATCTCGGAGA GGGAGGAGAA ATTCTTAAAA ACAGCTGCTC TCCTGCGCCC 1301 AGCTTCACAG GCAGCCCTGC CCCTTTCTCC TCACCAGCAT GGTACCTGCC 1351 CTTACTGCTA GAGCAGCTGC TTGTAGAGGG ACATTCCCTC CTTCCCAGTT TTAACTGGTG GACCACAGTG GGGGGAAAAA CATTCAAGCG ATATAAAGAC 1401 ACTTGGGCTC TTTGCAGATG CCTATACTTC CAACACTACC ATGTCCACAA 1451 ACCACCTGG GGGAGGCCC TTCCAAAGGG AGGCTTGCTA GTTTCAGCGT 1501 1551 CTAGCAGTTG GGTCCTCACT TTTACTCCAA TTGTGAAAAT AGCCCACGTA 1601 CCCTCGCAGT GTCCAGTAGG GATCCCAGAG GCACATAACC AAGAAAGGAT TTTGACTTTG TCACAGTGAC TATTTAAAAT AATCTATTCG AAGTCCAAAC 1651 1701 CAAACACAA GCCTGTGATA TTTTAGGTTA TTAAGGTAAC TGCTAATGAA 1751 GGATTTTAAA AAGTGTCCTC AAAAAGGACT TAGCCCCGGG AGTTGTTTAT 1801 AAAATTTCCC CCACTTGTAT ACAGTGTGCA CTAAAAGAAA ATGTATTTTA 1851 ATATCTAATG CCTGGGCTCT GAGCGTCATG CTTCTTGGTG GTAAACATGC 1901 AGTCCTGTTC CTAAGTGACT CAGAGGCATC AGAATTTCTC CACGTTACCC 1951 ATCTGCTTGG CACTCGGAAC TGAGCGTGTG AAATCCATAG CGCTGCCCAC AACCTGTTCT CACTGCTTAG CTCCCAGCTG GATTAAAGAC ACCTGCTCAG 2001 GCGGGAGAGA GAGAGAGAGA GCGAGCTTTT ACCTTGGAAA AGGTAAAGAT 2051 2101 GGAAATGTAC ACCAAAAAG ACAATTTTTA CATTTAATGG AACATTCTTT TTTTTTACAA GTATATTTTT CTACTGATAG TTTCAGAACA CTAATCTTAT 2151 ATTCACTCTA ATCTTAAACA TGTTTCTTTA AATATTTATA AGGCAGTTTA 2201 TTACAGAATA TTTTCATGCA ATCATGTGCA CATTATTGGT AGCAAACATA 2251 2301 GTATATCCTT TAGTACTTTA GCATATTTTT GTTAAAATAC TTTTAATGGT 2351 AAGAATGAA CTTGAGGTCC CAGGAGGTTT TGTTGCCTTT TCATTGATTA 2401 GAGACAATAA ATATCTTGTA ACTTCCTAAC CAGATCTGAG CTGTGCTCAC 2451 AATAATAATA ATGAAATCAG ATTCTTTGAT GCTGGACTCA GGAGGGAAAT

3751

CATTAGCCAA CTGTTGACTT ACTTATAGCT AGATGTCTAT GTGAGAAAGT 2501 ATAATATATA TATATACACA TATATATGAC ATGTAAGAGT CACTTTTATT 2551 TATCTGTCTT TGTTCACTTA TGAAGCCGGT AACTGCAGCA GTATGTTGGT 2601 GATGTCATGA TGCACAGAAG TCCCATGTGG AGTGTTTTTC CCACACTGAC 2651 AACTTGGCCT CCTTTCTGTG TGTTCAGTCT GTTGTCTGAA CTAACACTCC 2701 CGCGAGCACT ATACTCTTTA TACTCTGATC CCCCTAGTTC ATCTTAAATT 2751 TGTCTGTGGC CCTGGCAAGA TAGCGTACAC AAGATTCCAT GACTCCAGAG 2801 CATCTTGAAG AAACATACAT ATTTTGAAAG AGGGGAAATG TAGCAGATAG 2851 TTCACAAGCT GCGGGTTGTA GCTAAATATT CCATTTCTTT GAAATCATGT 2901 TTCTAAATTC TTTACCATCA GAAAGAAAAG GAGTGTCATA CACTTTCAAG 2951 GGAAGGCTTG GTCTGCGTTT TCTGTGTTTG GATTATTTT ATACTTTGCT 3001 GATCTTTAAG CTATCCATGG GGGAAATTTT ATACCAACGA GTTAATAATT 3051 CTCATTCATC GTTTACACAA TGTAACATGT GTCATACTGG GGCCAGCGAG 3101 ATGGCTCAGT AGGTAAAGGT GCTTGATGCT AAGCCCGGCA GCCTGTGTTT 3151 3201 CATCTACAGG ATGCACAACA TAAAAGAAAA GATCTGATTC CCACAGGTTC TCTTCTGACC TACACACAC CACACTAAAA TAACATTTAA AAATATGTGC 3251 CAAATTATAT TTGTTCGGGT GCCACCTTCC ACCAGCTTAC CACTACGGTA 3301 3351 GAACTGTCAA ATTCATCTCC CTGAATTTGT CTTAAAGGGG TGTCCATGCA CAGGCCCAAG AGTCACCTCC AATGAAATAA ATGTAATACT GAAGTATGCC 3401 ATGATGTTTG TTGTTTTCTT TCATCGTAAG CCTGTAAGCA GGAAAAATAC 3451 GTCAAATCAG ATAGAATAGA GCATTTACCA GTGGTCGATG GCCTGGTCAG 3501 TCCTGTGCCG GGTGACTTAG GACCAGGCAC GTCAGCTCTC TGAGCCTCCC 3551 CTTCCCTTGT TGTCACAAGG GAATAGAAGC AGAAGAAGCT GAGAGCCTCC 3601 CTATTCCCAG ATGCCCTGGT GGAATGACCT GCCTCTCTGC CGTTTCTGCC 3651 AACGTGTTGG TGCTATAAGC TGCTTCAAAT ACCAGTTTGT CTGTAGTGTG 3701 TACTCACCTA ATCACTTGTT ATCCAGTGCC TGTTCTAGGT TTATGGACTT

3801	AACTATTTCT	GTGATGTTTC	ATTTTTAGCC	ATGTTAACTC	CTAACACATA
3851	TTCTCTTATG	TCTCAGTAAA	GTTTCATTTG	ATAAGTTGTT	GAGATTCTGT
3901	TATTTGATAA	TATTCTTCGG	CTGTCCATCC	AGCATCTTAA	TCACTTTAAA
3951	ACTGTGATTG	TTATTTGCAA	CTCTGTTCTT	TGGAAAGAAT	AAAAGCATTT
4001	TTTTTCACTT	GCTAACATGC	TCACAAATGT	GAGAGAAGAG	TCATTAAAAG
4051	CTTTACTTAC	TGGGTCAGTG	CGTCATTGAC	TCCTTTCTGT	GTTTTGCCCA
4101	АТАААТТААТ	AAAAGACCAA	AAAAAAAAA	AAAAAAAAA	AAAAAA

Figure 3: cDNA sequence of mGy12 variant 2 (SEQ ID NO:3)

1	GCAGCGGCGG	CGGCGGCGAC	GACGGCGAAG	AGTTCATATG	GGAAGGACTT
51	TGGGGTGAGC	ATCTTCTCTA	TTTCCAGCTG	GCTTTTCTGA	TTCACCCCAC
101	CATTTAAAAC	CTGGAGGCAC	TGGGCCACAC	AAAGCCTTGC	TGATTTTCAG
151	AAAGAAGACT	CATCAAAGAT	GTCCAGCAAG	ACGGCAAGCA	CCAACAGCAT
201	AGCCCAAGCC	AGGAGAACTG	TGCAGCAGCT	GAGATTGGAA	GCCTCCATCG
251	AAAGAATAAA	GGTCTCAAAA	GCATCAGCAG	ACCTGATGTC	ATACTGTGAG
301	GAGCATGCCC	GGAGCGACCC	CCTGCTGATG	GGCATACCGA	CCTCAGAAAA
351	CCCGTTCAAG	GATAAGAAGA	CCTGCATCAT	CTTATAGTGG	ACCAGGAAGC
401	GCCCCTTGCC	TCTTAACGCA	AACCACAGCA	GCAACCTGAA	GGGATTCCTT
451	CAGCTTACCT	GGTAACCACA	GCTAGTAACT	AAAACACCCT	TCTCTCGGAA
501	TAATAGACCC	TGAAGTCTCT	CTTTTTCAAG	TTGTCCTTTC	TTCACCCTTT
551	ACTGATTTAA	TACAGAATAA	CAATCTTATT	TTCTATTTGA	TAACTATGGT
601	ATCATATTGG	GTTACTGTAT	AAGGAAAATG	GCAGGGGAGT	TGTGGGAAGC
651	TTGTCTTTAC	AAAATATAAT	TGATTAAGAT	ATGTCAAGAC	CTACATTGTC
701	TAAGCACCGG	CAAATTAAAA	TGTCGAGAAT	CACTTCAGTC	AAAAACCTTT
751	ATATTCTGTT	CTTAATAATG	TTTGTGCCAA	CCTATATCCC	ATGTAAGGGA
801	TCTGGGGAGG	AGGCATGTGT	CTACAACCAT	ACCTTTTTGC	ACTATGGGCA
851	CTAACCACCC	TGAAACTTCC	TGCGGTAGCT	CCCTCCCTTC	AGAGTTACAT
901	CATTATCCTG	ACTCTGTGTA	GGTAAATTTC	CGTGAAATTT	TTGTACAAAA
951	AAAAGGTAAT	GAAAGAACGT	TGCAAAGATC	ATCTGCATTA	TAATGAGTTG
1001	ATGCTGTTCT	CACTCCTCTC	TTGGAATTGT	GCTGGCCCCT	TAGTCTACAA
1051	TAAACTGTGC	CAATTAAAAA	CCTAAGGCTA	AAACTGAAAG	CCCTTTGATG
1101	GGGTCTTAAC	TCATATCAGT	CATTTGGGCT	TCTCTGATCC	TGAGGCTAAG
1151	AAAGGGGAAG	AGACCCTCAG	GAGGCAGCTT	CCACTCCAGG	GCTCTTGATC

1201 TCTGCTGGAT TGGGGGTGGC CACCTCAGAA ACTTCCACCC TCATGACTGG AATGGAAGAG GGGACCGAGA GCCTCACAAT CTCGGAGAGG GAGGAGAAAT 1251 TCTTAAAAAC AGCTGCTCTC CTGCGCCCAG CTTCACAGGC AGCCCTGCCC 1301 CTTTCTCCTC ACCAGCATGG TACCTGCCCT TACTGCTAGA GCAGCTGCTT 1351 GTAGAGGGAC ATTCCCTCCT TCCCAGTTTT AACTGGTGGA CCACAGTGGG 1401 1451 GGGAAAACA TTCAAGCGAT ATAAAGACAC TTGGGCTCTT TGCAGATGCC 1501 TATACTTCCA ACACTACCAT GTCCACAAAC CACCCTGGGG GAGGGCCCTT 1551 CCAAAGGGAG GCTTGCTAGT TTCAGCGTCT AGCAGTTGGG TCCTCACTTT TACTCCAATT GTGAAAATAG CCCACGTACC CTCGCAGTGT CCAGTAGGGA 1601 TCCCAGAGGC ACATAACCAA GAAAGGATTT TGACTTTGTC ACAGTGACTA 1651 TTTAAAATAA TCTATTCGAA GTCCAAACCA AACACAAAGC CTGTGATATT 1701 TTAGGTTATT AAGGTAACTG CTAATGAAGG ATTTTAAAAA GTGTCCTCAA 1751 1801 AAAGGACTTA GCCCCGGGAG TTGTTTATAA AATTTCCCCC ACTTGTATAC 1851 AGTGTGCACT AAAAGAAAAT GTATTTTAAT ATCTAATGCC TGGGCTCTGA 1901 GCGTCATGCT TCTTGGTGGT AAACATGCAG TCCTGTTCCT AAGTGACTCA 1951 GAGGCATCAG AATTTCTCCA CGTTACCCAT CTGCTTGGCA CTCGGAACTG 2001 AGCGTGTGAA ATCCATAGCG CTGCCCACAA CCTGTTCTCA CTGCTTAGCT CCCAGCTGGA TTAAAGACAC CTGCTCAGGC GGGAGAGAGA GAGAGAGAGC 2051 GAGCTTTTAC CTTGGAAAAG GTAAAGATGG AAATGTACAC CAAAAAAAGAC 2101 AATTTTACA TTTAATGGAA CATTCTTTTT TTTTACAAGT ATATTTTCT 2151 ACTGATAGTT TCAGAACACT AATCTTATAT TCACTCTAAT CTTAAACATG 2201 TTTCTTTAAA TATTTATAAG GCAGTTTATT ACAGAATATT TTCATGCAAT 2251 CATGTGCACA TTATTGGTAG CAAACATAGT ATATCCTTTA GTACTTTAGC 2301 2351 ATATTTTGT TAAAATACTT TTAATGGTAA GAAATGAACT TGAGGTCCCA 2401 GGAGGTTTTG TTGCCTTTTC ATTGATTAGA GACAATAAAT ATCTTGTAAC

3701

TTCCTAACCA GATCTGAGCT GTGCTCACAA TAATAATAAT GAAATCAGAT 2451 TCTTTGATGC TGGACTCAGG AGGGAAATCA TTAGCCAACT GTTGACTTAC 2501 TTATAGCTAG ATGTCTATGT GAGAAAGTAT AATATATAT TATACACATA 2551 TATATGACAT GTAAGAGTCA CTTTTATTTA TCTGTCTTTG TTCACTTATG 2601 AAGCCGGTAA CTGCAGCAGT ATGTTGGTGA TGTCATGATG CACAGAAGTC 2651 CCATGTGGAG TGTTTTCCC ACACTGACAA CTTGGCCTCC TTTCTGTGTG 2701 TTCAGTCTGT TGTCTGAACT AACACTCCCG CGAGCACTAT ACTCTTTATA 2751 CTCTGATCCC CCTAGTTCAT CTTAAATTTG TCTGTGGCCC TGGCAAGATA 2801 GCGTACACAA GATTCCATGA CTCCAGAGCA TCTTGAAGAA ACATACATAT 2851 TTTGAAAGAG GGGAAATGTA GCAGATAGTT CACAAGCTGC GGGTTGTAGC 2901 TAAATATTCC ATTTCTTTGA AATCATGTTT CTAAATTCTT TACCATCAGA 2951 AAGAAAAGGA GTGTCATACA CTTTCAAGGG AAGGCTTGGT CTGCGTTTTC 3001 TGTGTTTGGA TTATTTTAT ACTTTGCTGA TCTTTAAGCT ATCCATGGGG 3051 GAAATTTTAT ACCAACGAGT TAATAATTCT CATTCATCGT TTACACAATG 3101 TAACATGTGT CATACTGGGG CCAGCGAGAT GGCTCAGTAG GTAAAGGTGC 3151 TTGATGCTAA GCCCGGCAGC CTGTGTTTCA TCTACAGGAT GCACAACATA 3201 AAAGAAAAGA TCTGATTCCC ACAGGTTCTC TTCTGACCTA CACACACACA 3251 CACTAAAATA ACATTTAAAA ATATGTGCCA AATTATATTT GTTCGGGTGC 3301 CACCTTCCAC CAGCTTACCA CTACGGTAGA ACTGTCAAAT TCATCTCCCT 3351 GAATTTGTCT TAAAGGGGTG TCCATGCACA GGCCCAAGAG TCACCTCCAA 3401 TGAAATAAAT GTAATACTGA AGTATGCCAT GATGTTTGTT GTTTTCTTTC 3451 ATCGTAAGCC TGTAAGCAGG AAAAATACGT CAAATCAGAT AGAATAGAGC 3501 ATTTACCAGT GGTCGATGGC CTGGTCAGTC CTGTGCCGGG TGACTTAGGA 3551 CCAGGCACGT CAGCTCTCTG AGCCTCCCCT TCCCTTGTTG TCACAAGGGA 3601 ATAGAAGCAG AAGAAGCTGA GAGCCTCCCT ATTCCCAGAT GCCCTGGTGG 3651 AATGACCTGC CTCTCTGCCG TTTCTGCCAA CGTGTTGGTG CTATAAGCTG

3751	CTTCAAATAC	CAGTTTGTCT	GTAGTGTGTA	CTCACCTAAT	CACTTGTTAT
3801	CCAGTGCCTG	TTCTAGGTTT	ATGGACTTAA	CTATTTCTGT	GATGTTTCAT
3851	TTTTAGCCAT	GTTAACTCCT	AACACATATT	CTCTTATGTC	TCAGTAAAGT
3901	TTCATTTGAT	AAGTTGTTGA	GATTCTGTTA	TTTGATAATA	TTCTTCGGCT
3951	GTCCATCCAG	CATCTTAATC	ACTTTAAAAC	TGTGATTGTT	ATTTGCAACI
4001	CTGTTCTTTG	GAAAGAATAA	AAGCATTTTT	TTTCACTTGC	TAACATGCTC
4051	ACAAATGTGA	GAGAAGAGTC	ATTAAAAGCT	TTACTTACTG	GGTCAGTGCG
4101	TCATTGACTC	CTTTCTGTGT	TTTGCCCAAT	AAATTAATAA	AAGACCAAAA
4151	AAAAAAAA	AAAAAAAAA	AAAA		

amino acid sequence of human Gy12 (SEQ ID NO:4) Figure 4:

- 1. MSSKTASTNN IAQARRTVQQ LRLEASIERI KVSKASADLM SYCEEHARSD 51. PLLIGIPTSE NPFKDKKTCI IL



FIGURE 5